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TABLE 1 Continued

	LEAARLKQKEQERKSLE-LEKQKED--AQRRV-QERDKQWLEHVQEEQP	669
	HKIKRDKLKEIERKRLEQIQKKLEDEAARKAKQGKENLWRESIRKEEEE	687
	RPRKPHEEDRLKREDSVRKKEAEERAKPEMQDKQSRLFHPHQEPAKLATO	
	KQK-----RLQEEKSQDKTQEEER-KAEA--KQSET-----	
SH3A	APWSTTEKGPLTISAQESVKVYYYRALYPFESRSHDEITIOPGDIVMVDE	769
	-----ASALVNYRALYPFEARNHDEMFSFGDIIOVDE	748
	SOTGEPGWLGGELKGKTGWFPANYAEKIPENEVPTPAKPVTDLTSAAPAK	
	KTVGEPGWLYGSFOGKFGWPCNYVEKVLSS-----KALSPK	
	LALRETPAPLPVTSSEPSTTPNNWADFSSTWPSSSNEKPETDNWDTWAAQ	869
	KALLPPTVSLSATST-SSQPPASVTDYHNV---SFSNLTVNTTWQ-----	827
	PSLTVPSAGQLRQRSAFTPATATGSSPSVPLGQGEKVEGLQAOALYPWRA	
	-----QKSAFTRTVSPGSV-SPIHGQGOAVENLKAOLCSWTA	
SH3B	KKDNHLNFNKSDVITVLEOODMWWFGEVOGOKGWFPKSYVKLISGPVRKS	969
	KKENHLNFSKHDVITVLEOENWWFGEVHGGRGWFPKSYVKLIPGNEVQR	914
	TSIDTGPTESPASLKRVASPAKPAIP-GEEFIAMYTESSEOGDLTFOO	
	GEPEALYAAVTKKPTSTAYPVTSTAYPVGEDYIALYSYSSVEPGDLTFTE	
SH3C	GDVIVVTKKDGDWWTGTVGDKSGVFPSNYVRLKDSESGTAGKTGSLGKK	1068
	GEEILVTOKDGEWWTGSIGERTGIFPSNYVRPKDQENFGNASKSGASNKK	1014
SH3D	PEIAQVIASYAATGPEOLTLAPGOLILIRKKNPGGWEGELOARGKKROT	
	PEIAOVTSAYAASGTEOLSLAPGOLILILKNTSGWWOGELOARGKKROK	
	GWFPANYVKLLSPGTSKITPTLPTAVQPAVCQVIGMYDYTAONDDLA	
	GWFPASHVKLLGPSSERTMPT-----FHAVCQVIAMYDYMANNEDLN	
SH3E	FSKGOIINVLNKEDPDWWKGEVSGOVGLFPSNYVKLTTDMDPSQQ	1213
	FSKGOLINVMNKDDPDWOGETNGLTGLFPSNYVKMTTSDPSQQWCADL	1157
	QALDTMQPTERKRQGYIHELIOQTEERYMDDLQLFEQKTL	1197